

RAW SEQUENCE LISTING

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Application Serial Number: 10/549,262
Source: PCT
Date Processed by STIC: 09/28/2005

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RAW SEQUENCE LISTING

DATE: 09/28/2005

PATENT APPLICATION: US/10/549,262

TIME: 13:52:32

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\09282005\J549262.raw

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3 <110> APPLICANT: Forschungszentrum Jlich GmbH
5 <120> TITLE OF INVENTION: Method for Microbial Production of L-Serine
7 <130> FILE REFERENCE: 23369
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/549,262
C--> 10 <141> CURRENT FILING DATE: 2005-09-12
12 <160> NUMBER OF SEQ ID NOS: 2
14 <170> SOFTWARE: PatentIn Ver. 2.1
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 1449
18 <212> TYPE: DNA
19 <213> ORGANISM: Corynebacterium glutamicum
21 <400> SEQUENCE: 1
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23 caccttgtag accaccatag aaaaggccca ccctcagcca tggctatcag tggtgttgat 120
24 ctatttagca tcggtatcgg accatcatcc tcacataccg tcggcccat gagagccgcc 180
25 ctcacgtata tctctgaatt tcccagctcg catgtcgata tcacgttgca cggatccctt 240
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27 gaaccaacga tagttcccat tgatgctgca ccctcaccgc gcgcgccgat tcctgcgaaa 360
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46 gagtgttag                                     1449
49 <210> SEQ ID NO: 2
50 <211> LENGTH: 449
51 <212> TYPE: PRT
52 <213> ORGANISM: Corynebacterium glutamicum
54 <400> SEQUENCE: 2
55 Met Ala Ile Ser Val Val Asp Leu Phe Ser Ile Gly Ile Gly Pro Ser

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56      1              5              10              15
58 Ser Ser His Thr Val Gly Pro Met Arg Ala Ala Leu Thr Tyr Ile Ser
59              20              25              30
61 Glu Phe Pro Ser Ser His Val Asp Ile Thr Leu His Gly Ser Leu Ala
62              35              40              45
64 Ala Thr Gly Lys Gly His Cys Thr Asp Arg Ala Val Leu Leu Gly Leu
65              50              55              60
67 Val Gly Trp Glu Pro Thr Ile Val Pro Ile Asp Ala Ala Pro Ser Pro
68 65              70              75              80
70 Gly Ala Pro Ile Pro Ala Lys Gly Ser Val Asn Gly Pro Lys Gly Thr
71              85              90              95
73 Val Ser Tyr Ser Leu Thr Phe Asp Pro His Pro Leu Pro Glu His Pro
74              100             105             110
76 Asn Ala Val Thr Phe Lys Gly Ser Thr Thr Arg Thr Tyr Leu Ser Val
77              115             120             125
79 Gly Gly Gly Phe Ile Met Thr Leu Glu Asp Phe Arg Lys Leu Asp Asp
80              130             135             140
82 Ile Gly Ser Gly Val Ser Thr Ile His Pro Glu Ala Glu Val Pro Cys
83 145             150             155             160
85 Pro Phe Gln Lys Ser Ser Gln Leu Leu Ala Tyr Gly Arg Asp Phe Ala
86              165             170             175
88 Glu Val Met Lys Asp Asn Glu Arg Leu Ile His Gly Asp Leu Gly Thr
89              180             185             190
91 Val Asp Ala His Leu Asp Arg Val Trp Gln Ile Met Gln Glu Cys Val
92              195             200             205
94 Ala Gln Gly Ile Ala Thr Pro Gly Ile Leu Pro Gly Gly Leu Asn Val
95              210             215             220
97 Gln Arg Arg Ala Pro Gln Val His Ala Leu Ile Ser Asn Gly Asp Thr
98 225             230             235             240
100 Cys Glu Leu Gly Ala Asp Leu Asp Ala Val Glu Trp Val Asn Leu Tyr
101              245             250             255
103 Ala Leu Ala Val Asn Glu Glu Asn Ala Ala Gly Gly Arg Val Val Thr
104              260             265             270
106 Ala Pro Thr Asn Gly Ala Ala Gly Ile Ile Pro Ala Val Met His Tyr
107              275             280             285
109 Ala Arg Asp Phe Leu Thr Gly Phe Gly Ala Glu Gln Ala Arg Thr Phe
110              290             295             300
112 Leu Tyr Thr Ala Gly Ala Val Gly Ile Ile Ile Lys Glu Asn Ala Ser
113 305             310             315             320
115 Ile Ser Gly Ala Glu Val Gly Cys Gln Gly Glu Val Gly Ser Ala Ser
116              325             330             335
118 Ala Met Ala Ala Ala Gly Leu Cys Ala Val Leu Gly Gly Ser Pro Gln
119              340             345             350
121 Gln Val Glu Asn Ala Ala Glu Ile Ala Leu Glu His Asn Leu Gly Leu
122              355             360             365
124 Thr Cys Asp Pro Val Gly Gly Leu Val Gln Ile Pro Cys Ile Glu Arg
125              370             375             380
127 Asn Ala Ile Ala Ala Met Lys Ser Ile Asn Ala Ala Arg Leu Ala Arg
128 385             390             395             400

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```
130 Ile Gly Asp Gly Asn Asn Arg Val Ser Leu Asp Asp Val Val Val Thr
131                               405                      410                      415
133 Met Ala Ala Thr Gly Arg Asp Met Leu Thr Lys Tyr Lys Glu Thr Ser
134                               420                      425                      430
136 Leu Gly Gly Leu Ala Thr Thr Leu Gly Phe Pro Val Ser Met Thr Glu
137                               435                      440                      445
139 Cys
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VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Application Number

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date